

Validation of Bioassays for Vaccines

Bob Dillard

Takeda Pharmaceuticals North America

Presented at the NIH sponsored meeting

Assaying Potency of Novel Vaccines

October 11, 2005

Outline

- Why validate?
- Assay characteristics → Model
- Assay validation parameters
- Reportable value → Power of averaging
- Acceptance criteria

Why Validate?

- Regulatory Expectations
 - Measurement is the foundation on which research decisions rest
 - Don't think of validation as pass/fail
 - Use the validation results to inform your routine choices
- ➔ replication – informed averaging

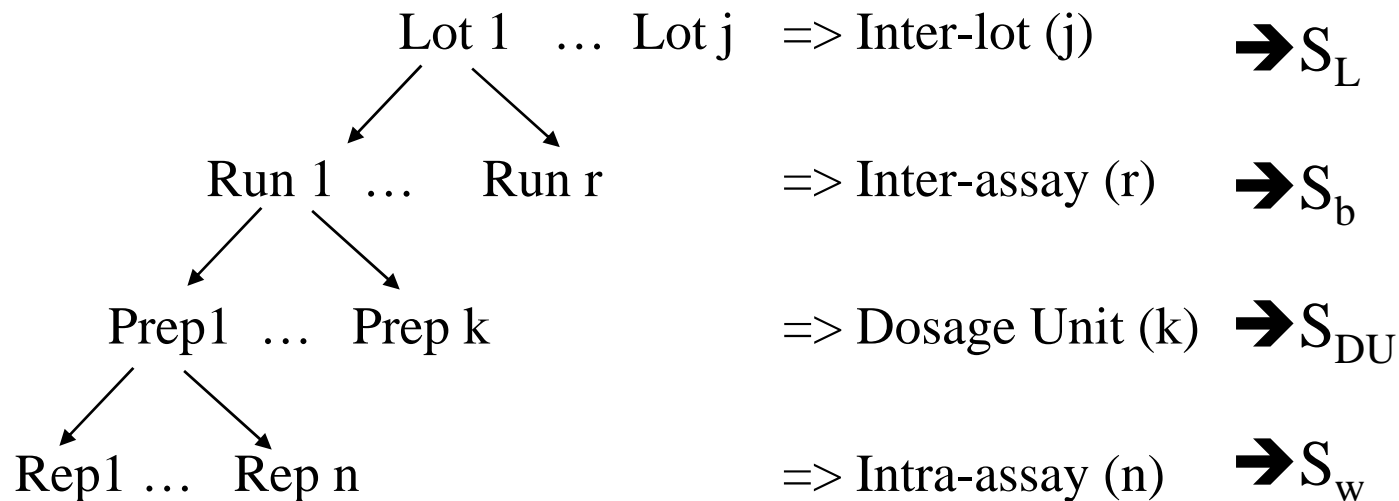
When ?

- As soon as you start making decisions using the data
- After the optimal operating conditions have been established in assay development – stable operating conditions
 - Driving intra-assay factors
 - Driving Inter-assay factors
- Continuous assessment

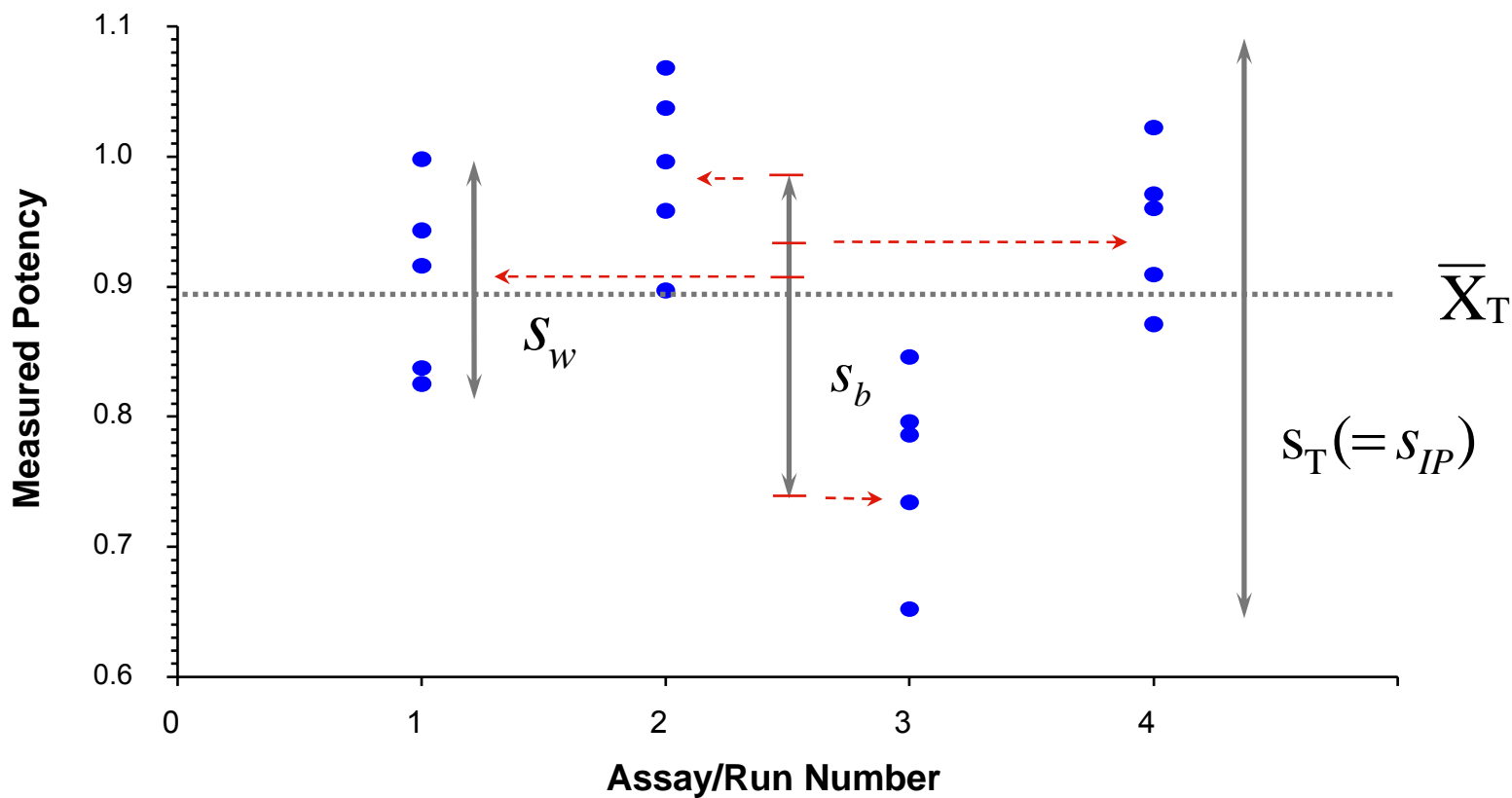
➔ Early and continuously thereafter

Assumptions

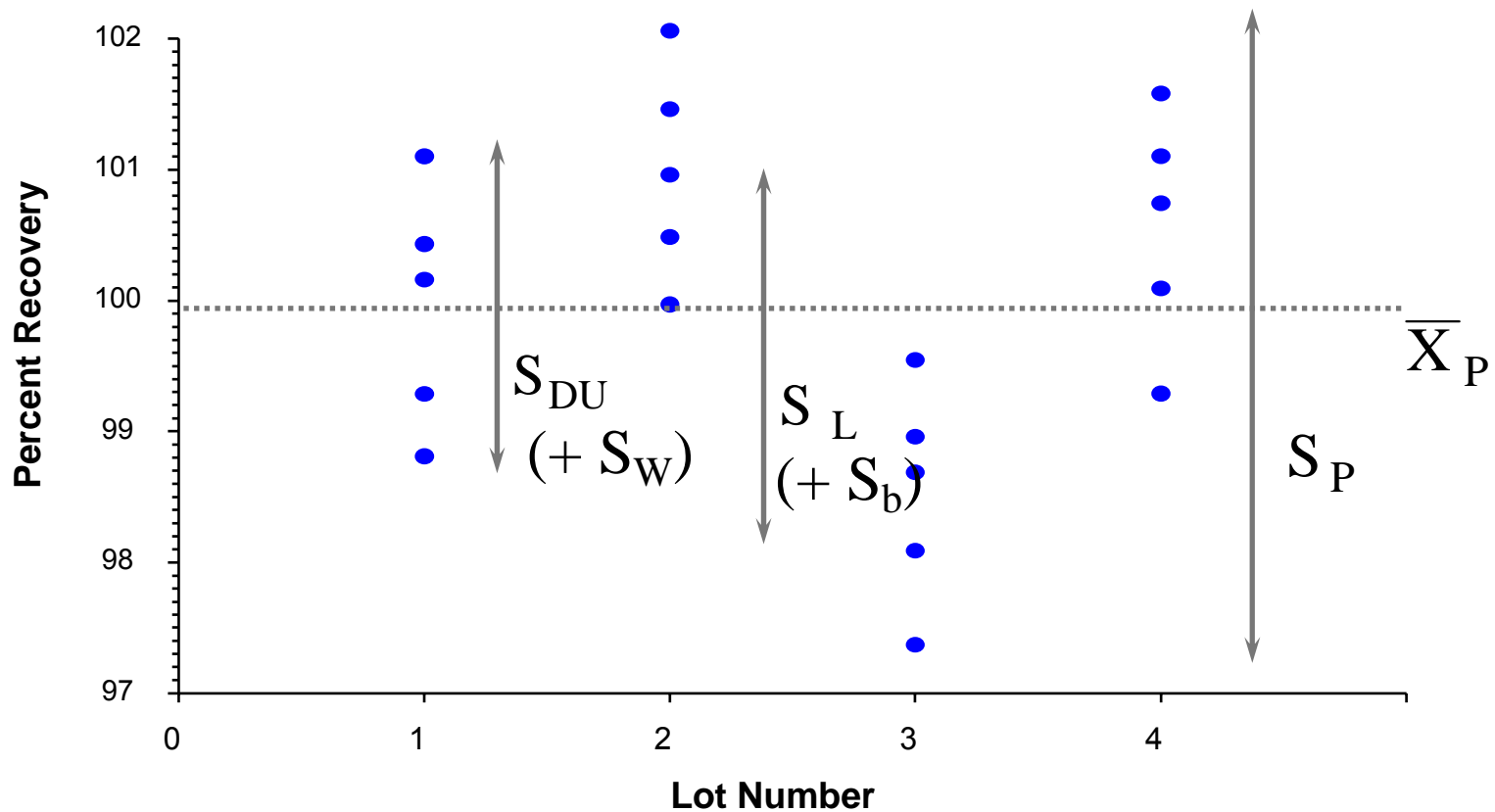
- Measure is a biologic response (activity not mass)
 ➔ highly variable
- Continuous response – or at least convertible
- Simplistic Layout



Model of the Assay



Broader Model – Process (& Assay)



Assay Validation Parameters

■ Relative “accuracy”/linearity

- Dilution effect
 - Forced degradation
- $$\left. \vphantom{\begin{matrix} \text{Dilution effect} \\ \text{Forced degradation} \end{matrix}} \right\} \overline{X}_A + \overline{X}_P$$

■ Precision

- Robustness – intra-assay factors
- Ruggedness – inter-assay factors
- **Reproducibility** – random draw routine control - $S_T(\rightarrow S_{RV})$

■ Others are variations on accuracy or precision

- Limit of detection, range, interference

Validation Design - Precision

- Replication pattern targeted toward primary noise sources
 - Intra-run noise
 - Inter-run noise
 - Run is independent preparation of reagents, test, and standards
 - Don't short change the number of runs (≥ 6)
- Choice of levels dictated by range of product potency
- Avoid pass/fail mentality – worst case, not best

Assay Capability – Getting the Numbers

- Estimates based on standard deviations of the individual replicates and the run averages.

$$\left. \begin{aligned} s_w &= \text{Avg}(s_{Run1}, s_{Run2}, \dots) \\ s_b &= s_{\bar{X}} - (s_w / n) \end{aligned} \right\} \text{ANOVA}$$

- 3 Rules:

- Noise is cumulative $\rightarrow S_{All}^2 = S_1^2 + S_2^2 + \dots$
- Averaging improves precision – **predictably** $\rightarrow S_{Avg}^2 = \frac{S_{y_i}^2}{n}$
- Mean $\pm 3s$ bracket the results

Assay Capability - Using the Information

■ Reportable Value

- What constitutes an assay?

■ Sources of Noise (Propagation of Errors)

- Control by averaging

$$S_{RV}^2 = \frac{S_{DU}^2}{k} + \frac{S_W^2}{n} + S_b^2$$

Average of c composites:

$$S_{RV}^2 = \frac{S_{DU}^2}{c * k} + \frac{S_W^2}{c * n} + S_b^2$$

/ r

In viral/bioassay average over r runs

Establishing a Reportable Value

- How do you define the rv?

➔ Impact, Criteria, Cost

- Impact of the sample allocation.

$$\text{Suppose, } \left. \begin{array}{l} s_w = 17\% \\ s_b = 25\% \end{array} \right\} S_{RV} = \sqrt{\frac{S_w^2}{n \cdot r} + \frac{S_b^2}{r}}$$

n	Runs (r)		
	1	6	12
1	30%	12%	9%
2	28%	11%	8%
3	27%	11%	8%

Acceptance Criteria

- Acceptance Criteria dictated by use of the assay
 - Define use by a range or specification limits
- Adjust the replication so that,

$$6\sqrt{S_P^2 + S_{RV}^2} \leq Range$$

- If our desired range is 50% to 150%
→ $S_{RV} \leq 17\%$

How much replication is too much?

■ Replication vs Method Improvement

- Partly driven by \$

■ Capability of the Art ?

	S_T	Range
• Small Molecule (HPLC)	<5%	±15%
• SM in matrix (GC/Mass Spec)	15%	±45%
• Bigger Molecule (Immunoassay)	20%	±60%
• BM activity (Bioassay)	50%	±150%
• Viral Assay		

But these can be easily reduced by 60% just by judicious averaging

So, what level of S_{RV} do I target?

Pitfalls

■ Limited data has risks

- Some risks are controlled by choice of multipliers
- Look for ways to update and expand information

➔ Follow-up (continuous assessment)

Stability studies, Control samples, Scale-up

■ The curse of the validation experiment

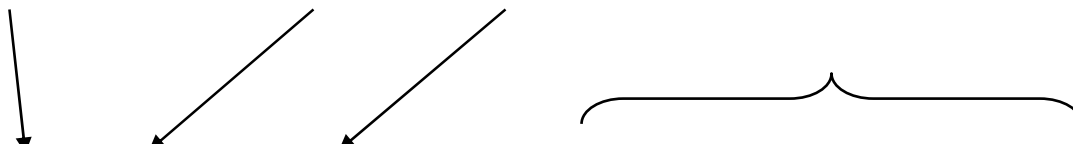
- We tend to reward pass/fail rather than good information

■ You will likely need to work in log scale

Data Driven Release Specifications

Data Driven Expiry Specifications

ES = Process/Assay Mean \pm Drift – Degradation \pm Uncertainties

$$L/UES = \bar{X} \pm S_L + \beta * T \pm 3 * \sqrt{S_L^2 + T^2 * S_\beta^2 + S_{RV}^2}$$


Then ask:

Are these specs pharmacologically sound?

Are they close to what agencies are asking for?

Recommendations

- Define carefully what values should be held up to the specification – reportable value.
 - Do NOT expect individual values to meet those same specs.
 - Paradox of individuals – disincentive to collect more data.
- Validation is a continuous process
 - Utilize all of your information